

REPLACEMENT SHEET

Fig. 15A

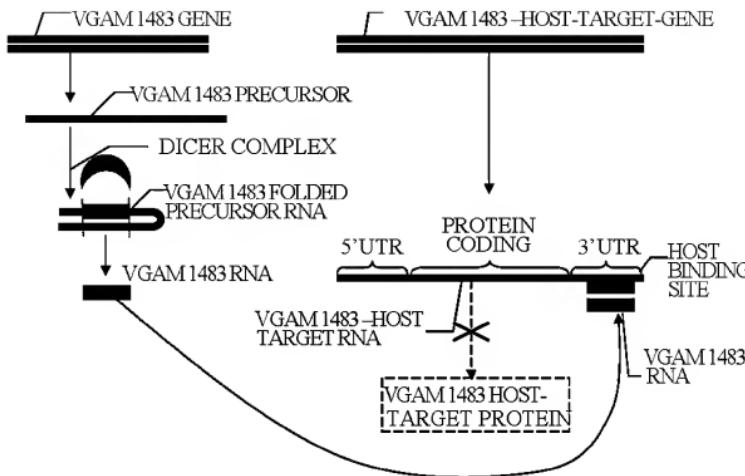


Fig. 15B

GTGTGACGTGAGTCTGATCCAACACTGAACGCTTCGTC SEQ ID: 37404

GTGTTTTCATGCAGTTTACAGACCATGACAAGCCTGA

CGAGAGCGTTCATCGGGCATGAAGTACGCATTACAC

TGACAAGCCTGACGAGAGCGTTCA

SEQ ID: 37405

Fig. 15C

—	AG	—	A	AACAC	T	T	CAG	TT	
GTGTGA	CGTG	TC	TG	TCC	TGAAACGCTTCGTCG	GTTT	TCATG	CT	T
CACATT	GCAT	AG	AC	GGG	<u>ACTTGCGAGAGCAGT</u>	CGAA	AGTAC	GA	/
AC	GA	T	—	GCT—	C	C	CA	—	CA

REPLACEMENT SHEET

Fig. 15D/1

	CAA	CGA		
C7	5' TGA	GCCTGA	GAGCGTTCA	3'
BINDING SITE				
	3' ACT	CGGACT	CTCGTAAGT	5'
	ATC	TAC		
	A	ACGA	G	
DUSP5	5' TG	CAAGCCTG	GA	CGTTCA
BINDING SITE				
	3' AC	GTTCGGAC	CT	GTGAGT
	-	CA--	G	
	A	AC		
ATP6V1F	5' TG	CAAGCCTG	GAGAG	3'
BINDING SITE				
	3' AC	GTTCGGAC	CTCTC	5'
	C	CC		
	CAA	AGAGCG		
DKFZP564C103	5' TGA	GCCTGAGC	TTCA	3'
BINDING SITE				
	3' ACT	CGGACTGC	AAGT	5'
	AC-	GAGAG-		
	CA	C	C	
KIAA0561	5' GA	AGC	TGACGAGAG	GT
BINDING SITE				
	3' CT	TCG	ACTGCTCTC	CA
	AC	A	C	
	A	A G	A	
KIAA0935	5' TG	CAAGCCTG	C	AG
BINDING SITE				
	3' AC	GTTCGGAC	G	TC
	C	C	C	
	AGC			
MIC2L1	5' GACAAGCCTGACGAG		GTTCA	3'
BINDING SITE				
	3' CTGTTCGGACTGTT		CAAGT	5'

	C	C-		
PPP4R1L	5' TGACAAG	CTGA	GAGAGCG	3'
BINDING SITE				
	3' ACTGTT	GACT	TTCTCGC	5'
	A	AC		
	TGACG	GC-		
ZAK	5' TGACAAGCC	AGA	GTTCA	3'
BINDING SITE				
	3' ACTGTTCGG	TCT	CAAG	5'
	-----	AGT		

REPLACEMENT SHEET

Fig. 15D/2

	CC	ACGA					
LOC149566	5' TGACAAG	TG	GAGCGTT	3'	SEQ ID:37405		
BINDING SITE							
	3' ACTGTTCA	AC	CTTGCAA	5'	SEQ ID:37415		
	A-	CCA-					
	A	AC	G	CG			
LOC150368	5' TG	CAAGCCTG	GA	AG	TTC	3'	SEQ ID:37405
BINDING SITE							
	3' AC	GTTCGGAC	CT	TC	GAG	5'	SEQ ID:37416
	C	A-	G	A-			
	A	CGAGA	C				
LOC255718	5' TGACA	GCCTGA	G	GTTC	3'	SEQ ID:37405	
BINDING SITE							
	3' ACTGT	CGGACT	C	CAGG	5'	SEQ ID:37417	
	C	-----	C				